PCT/AU00/00056

## WO 00/46Z33

1

Sequenc Listing:

<110>\The University of Sydney

<120> Pigment protein from coral tissue

<160> 15

<170> PatehtIn Ver. 2.1

<210> 1

<211> 5

<212> PRT

<213> Acropora aspera, Acropora horrida, Montipora caliculata, Porites murrayensis, Montipora monasteriata, and Porites lobata

<400> 1 Ser Val Ile Ala Lys

<210> 2

<211> 17

<212> PRT

<213> Acropora horrida

<400> 2

Ser Val Ile Ala Lys Gln Met Thr Tyr Lys Val Tyr Met Ser Gly Thr 1 15

Val

<210> 3

<211> 231

<212> PRT

<213> Acropora aspera

<400> 3

Ser Val Ile Ala Lys Gln Met Thr Tyr Lys Val Tyr Met Ser Gly Thr 1 5 15

Val Asn Gly His Tyr Phe Glu Val Glu Gly Asp Gly Lys Gly Lys Pro 20 25 30

Tyr Glu Gly Glu Gln Thr Val Arg Leu Ala val Thr Lys Gly Gly Pro

Leu Pro Phe Ala Trp Asp Ile Leu Ser Pro Gla Cys Gla Tyr Gly Ser

lle Pro Phe Thr Lys Tyr Pro Glu Asp Ile Pro Asp Tyr Val Lys Gln
65 70 75

Ser Phe Pro Gly Arg Tyr Thr Trp Glu Arg Ile Met Asn Phe Glu Asp

Gly Ala Val Cys Thr Val Ser Asn Asp Ser Ser Ild Gln Gly Asn Cys 100 110

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2

Phe Ile Tyr His Val Lys Phe Ser Gly Leu Asn Phe Pro Pro Asn Gly
115 - 125

Pro Val Med Gln Lys Lys Thr Gln Gly Trp Glu Pro Asn Thr Glu Arg

Leu Phe Ala Arg Asp Gly Met Leu Ile Gly Asn Asn Phe Met Ala Leu 145 150 155

Lys Leu Glu Gly Gly His Tyr Leu Cys Glu Phe Lys Ser Thr Tyr 165 170 175

Lys Ala Arg Lys Pro Val Lys Met Pro Gly Tyr His Tyr Val Asp Arg

Lys Leu Asp Val Thr Asn His Asn Lys Asp Tyr Thr Ser Val Glu Gln
195 200 205

Arg Glu Ile Ser Ile Ala Arg Lys Pro Leu Val Ala Cys Cys Phe Phe 210 220

Arg Val Lys Ser Arg His Lys 225 230

<210> 4

<211> 235

<212> PRT

<213> Acropora aspera

<400> 4
Ser Val Ile Ala Lye Gln Met Thr Tyr Lys Val Tyr Met Ser Gly Thr
1 10 15

Val Asn Gly His Tyr Phe Glu Val Glu Gly Asp Gly Lys Pro 20 25 30

Tyr Glu Gly Glu Gln Thr Val Arg Leu Ala Val Thr Lys Gly Gly Pro
35 40 45

Leu Pro Phe Ala Trp Asp Ile Leu Ser\Pro Gln Cys Gln Tyr Gly Ser
50 60

Ile Pro Phe Thr Lys Tyr Pro Glu Asp Ile Pro Asp Tyr Val Lys Gln 65 70 75 80

Ser Phe Pro Gly Arg Tyr Thr Trp Glu Arg\lle Met Asn Phe Glu Asp
85
90

Gly Ala Val Cys Thr Val Ser Asn Asp Ser Ser Ile Gln Gly Asn Cys
100 105 110

Phe Ile Tyr His Val Lys Phe Ser Gly Leu Ash Phe Pro Pro Asn Gly

Pro Val Met Gln Lys Lys Thr Gln Gly Trp Glu\Pro Asn Thr Glu Arg

Leu Phe Ala Arg Asp Gly Met Leu Ile Gly Asn Asn Phe Met Ala Leu 145 150 155 he het Ala Leu

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Lys Leu Glu Gly Gly Gly His Tyr Leu Cys Glu Phe Lys Ser Thr Tyr
Lys Ala Lys Lys\Pro Val Lys Met Pro Gly Tyr His Tyr Val Asp Arg
                                                            185
 Lys Leu Asp Val Thr Ash His Asn Lys Asp Tyr Thr Ser Val Glu Gln
                                                    200
 Cys Glu Ile Ser Ile Ala Arg Lys Pro Val Val Ala Cys Arg Phe Phe
                                            215
 Arg Val Lys Ser Arg Ris Lys Tyr Ala Val Ala
 225
                                      2)30
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 <211> 841
 <212> DNA
 <213> Acropora aspera
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 cagtacggaa gcataccatt caccalgtac cetgaagaca tecetgacta tgtaaagcag 240
tcattcccgg ggagatatac atgggagagg atcatgaact ttgaagatgg tgcagtgtg 300 actgtcagca atgattccag catccaaggc aactgtttca tttaacatgt caagttctct 360 ggtttgaact ttcccccaa tggacctgtt atgcagaaga agacacaggg ctgggaaccc 420 aacactgagg gtctctttgc acgagatgga atgctgatag gaaacaactt tatggctctg 480 aagttagaag gaggtggtca ctatttgfg gaattcaaat ctacttacaa ggcaaggaag 540 cctgtgaaga tgccagggta tcactatgtt gaccgcaaac tggatgtaac caatcacaac 600 aaggattaca cttccgttga gcagcgtgaa attcccattg cacgcaaacc tttggtcgcc 660 tgctgttttt tcagaggtga atcacagga aacacga tggcgtaaaa aacgtagatt 720
 tgctgttttt tcagagtcaa atcaaggcac aaataagcag tggcgtaaaa aacgtagatt 720
 ctgattttag cttagagaag taggaacgaa galgtgtaga caacettcaa tgattaaact 780 tttgaaaaca acsccaaaaa aaaaaaaaa aaaaaagcggc cgctcgaatt 840
 <210> 6
 <211> 841
 <212> DNA
 <213> Acropora aspera
 <400> 6
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cagtacggaa gcataccatt caccaagtac cotgaagaca tocotgacta tgtaaagcag 240 teattocogg ggagatatac atggagagag atcatgaact ttgaagatgg tgcagtgtgt 300 actgtcagca atgattccag catccaaggc aactgtttca totaccatgt caagttotot 360 ggtttgaact ttcetoccaa tggacctgtt atgcagaaga agacacaggg ctgggaacce 420
 aacactgage gtotetttge acgagatgga atgethatag gaaacaactt tatggetetg 480
 aagttagaag gaggtggtca ctatttgtgt gaattcaaat ctacttacaa ggcaaagaag 540 cctgtgaaga tgccagggta tcactatgtt gaccgcaaac tggatgtaac caatcacaac 600 aaggattaca cttccgttga gcagtgtgaa atttccattg cacgcaaacc tgtggtcgcc 600
 tgccgttttt tcagagtcaa atcaaggcac aaatacgcag tggcgtaaaa aacgtagatt 720
 ctgattttag cttatagaag taggaacgaa gaagtgtaaa caaccattaa tgattaaact 780
 tttgaaaaca acgccataaa aaaaaaaaaa aaaaaaaadaa aaaaagcggc cgctcgaatt 840
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<210> 7 <211> 18

<212> PRT <213> Acropora aspera, Montipora caliculata, and Porites murrayensis

Ser Val Ile Ala L\s Gln Met Thr Tyr Lys Val Tyr Met Ser Gly Thr

Val Asn

<210> 8

<211> 25 <212> PRT

<213> Porites lobata

Ser Val Ile Ala Lys Gln Med Thr Tyr Lys Val Tyr Met Ser Gly Thr

Val Asn Asn His Tyr Glu Phe Val Thr 20 25

<210> 9

<211> 225

<212> PRT

<213> Discosoma sp.

<400> 9

Met Arg Ser Ser Lys Asn Val Ile Lys Glu Phe Met Arg Phe Lys Val

Arg Met Glu Gly Thr Val Asn Gly His\Glu Phe Glu Ile Glu Gly Glu

Gly Glu Gly Arg Pro Tyr Glu Gly His Asn Thr Val Lys Leu Lys Val

Thr Lys Gly Gly Pro Leu Pro Phe Ala Tro Asp Ile Leu Ser Pro Gln

Phe Gln Tyr Gly Asn Lys Val Tyr Val Lys Wis Pro Ala Asp Ile Pro

Asp Tyr Lys Lys Leu Ser Phe Pro Glu Gly Phe Lys Trp Glu Arg Trp

Met Asn Phe Glu Asp Gly Gly Val Val Thr Val Thr Gln Asp Ser Ser

Leu Gln Asp Gly Cys Phe Ile Tyr Lys Val Lys Phe Ile Gly Val Asn

Phe Pro Ser Asp Gly Pro Val Met Gln Lys Lys The Met Gly Trp Glu 135 130

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5

Ala Ser Thi Lys Arg Leu Tyr Pro Arg Asp Gly Val Leu Lys Gly Glu 145 150 155 160

Ile His Lys Ala Leu Lys Leu Lys Asp Gly Gly His Tyr Leu Val Glu 165 170 175

Phc Lys Ser Ile Tyr Met Ala Lys Lys Pro Val Gln Leu Pro Gly Tyr 180 185 190

Tyr Tyr Val Asp Ser Lys Leu Asp Ile Thr Ser His Asn Glu Asp Tyr
195 200 205

Thr Ile Val Glu Glm Tyr Glu Arg Thr Glu Gly Arg His His Leu Phe 210 220

Leu 225

<210> 10

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<211> 230 <212> PRT

<213> Discosoma sp.

<400> 10

Met Ser Lys Gly Glu Glu Let Phe Thr Gly Val Val Pro Ile Leu Val 1 5 15

Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu 20 25 30

Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys

Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe

Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg

His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg

Thr Ile Phe Phe Lys Asp Asp Gly Ash Tyr Lys Thr Arg Ala Glu Val 100 105 110

Lys Phe Glu Gly Asp Thr Leu Val Ash Arg Ile Glu Leu Lys Gly Ile 115 125

Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn 130 140

Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly 145 155 160

Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val 165 170 175

Gln Leu Ala Asp His Tyr Gln Gln Asn The Pro Ile Gly Asp Gly Pro
180 185 190

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Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser

Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val 210 220

Thr Ala Ala Gly Ile Thr 225 230

<210>.11

WO 00/46233

<211> 20 <212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primers

<400> 11

teegttateg ctaaacagat

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<210> 12

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primers

<400> 12

tttgtgcctt gatttgactc

20

<210> 13

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primers

<400> 13

cgccactgcg tatttgtgcc

20

<210> 14

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR primers

<400> 14

ggcgaccaca ggtttgcgtg .

20

<210> 15

<211> 30

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<212> DNA <213> Artificial Sequence

<223> Description of Artificial Sequence: FCR primers

<400> 15 tccgttatcg ctaaacagat gacctacaaa

30

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